

Hilma Holm

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/6949461/publications.pdf>

Version: 2024-02-01

16
papers

1,103
citations

933264

10
h-index

940416

16
g-index

17
all docs

17
docs citations

17
times ranked

1305
citing authors

#	ARTICLE	IF	CITATIONS
1	The power of genetic diversity in genome-wide association studies of lipids. <i>Nature</i> , 2021, 600, 675-679.	13.7	353
2	Large-scale integration of the plasma proteome with genetics and disease. <i>Nature Genetics</i> , 2021, 53, 1712-1721.	9.4	340
3	Long-read sequencing of 3,622 Icelanders provides insight into the role of structural variants in human diseases and other traits. <i>Nature Genetics</i> , 2021, 53, 779-786.	9.4	156
4	A genome-wide meta-analysis yields 46 new loci associating with biomarkers of iron homeostasis. <i>Communications Biology</i> , 2021, 4, 156.	2.0	72
5	Distinction between the effects of parental and fetal genomes on fetal growth. <i>Nature Genetics</i> , 2021, 53, 1135-1142.	9.4	41
6	The genetic architecture of age-related hearing impairment revealed by genome-wide association analysis. <i>Communications Biology</i> , 2021, 4, 706.	2.0	30
7	Genetic insight into sick sinus syndrome. <i>European Heart Journal</i> , 2021, 42, 1959-1971.	1.0	27
8	MEPE loss-of-function variant associates with decreased bone mineral density and increased fracture risk. <i>Nature Communications</i> , 2020, 11, 4093.	5.8	24
9	Large-Scale Screening for Monogenic and Clinically Defined Familial Hypercholesterolemia in Iceland. <i>Arteriosclerosis, Thrombosis, and Vascular Biology</i> , 2021, 41, 2616-2628.	1.1	16
10	A genome-wide meta-analysis uncovers six sequence variants conferring risk of vertigo. <i>Communications Biology</i> , 2021, 4, 1148.	2.0	12
11	Predicting the probability of death using proteomics. <i>Communications Biology</i> , 2021, 4, 758.	2.0	10
12	Genetic variants associated with platelet count are predictive of human disease and physiological markers. <i>Communications Biology</i> , 2021, 4, 1132.	2.0	7
13	Population-level deficit of homozygosity unveils CPSF3 as an intellectual disability syndrome gene. <i>Nature Communications</i> , 2022, 13, 705.	5.8	7
14	Comment on "Evaluating the cardiovascular safety of sclerostin inhibition using evidence from meta-analysis of clinical trials and human genetics". <i>Science Translational Medicine</i> , 2021, 13, eabe8497.	5.8	4
15	Molecular benchmarks of a SARS-CoV-2 epidemic. <i>Nature Communications</i> , 2021, 12, 3633.	5.8	3
16	Genetic architecture of band neutrophil fraction in Iceland. <i>Communications Biology</i> , 2022, 5, .	2.0	1